

RAW SEQUENCE LISTING

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Application Serial Number: 10/524,564

Source: PG

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PCT

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DATE: 02/09/2006

PATENT APPLICATION: US/10/524,564

TIME: 12:26:02

Input Set : A:\26350095.APP

Output Set: N:\CRF4\02012006\J524564.raw

3 <110> APPLICANT: UEDA, HIROSHI
5 <120> TITLE OF INVENTION: A METHOD TO DETERMINE PROTEIN INTERACTIONS
7 <130> FILE REFERENCE: 026350-095
9 <140> CURRENT APPLICATION NUMBER: 10/524,564
10 <141> CURRENT FILING DATE: 2005-02-14
12 <150> PRIOR APPLICATION NUMBER: PCT/JP03/10386
13 <151> PRIOR FILING DATE: 2003-08-15
15 <150> PRIOR APPLICATION NUMBER: JP 2002-237411
16 <151> PRIOR FILING DATE: 2002-08-16
18 <160> NUMBER OF SEQ ID NOS: 41
20 <170> SOFTWARE: PatentIn Ver. 3.3
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 4301
24 <212> TYPE: DNA
25 <213> ORGANISM: Artificial Sequence
27 <220> FEATURE:
28 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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111 <220> FEATURE:
112 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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115 <400> SEQUENCE: 2
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196 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

197 primer for ompA secretion signal sequence

198 and FLAG tag sequence (Back: ompXbaRV)

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207 <213> ORGANISM: Artificial Sequence

209 <220> FEATURE:

210 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

211 primer for ompA secretion signal sequence and

212 FLAG tag sequence (Forward: ompApaSalFR)

214 <400> SEQUENCE: 4

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226     (Back:g7KpnRV)
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281     primer for split Fv linker (Back:LinkBackX)
283 <400> SEQUENCE: 9

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VERIFICATION SUMMARY

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